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_1: AA212	2542. mu4	14c08.r1 Soare	es[gi:1811196]			Links	
LOCUS DEFINITION			470 bp _thymus_2NbMT M			-FEB-1997 GE:642254	
ACCESSION VERSION KEYWORDS	EST.	42.1 GI:18					
SOURCE ORGANISM	Mus musculus (house mouse) Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.						
REFERENCE AUTHORS	1 (bases 1 to 470) Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T., Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and Waterston, R.						
TITLE JOURNAL COMMENT	The Wa Unpubl Contac WashU- Washin 4444 F Tel: 3 Fax: 3 Email:	shU-HHMI Mo ished (1996 t: Marra M/ HHMI Mouse gton Univer orest Park 14 286 1800 14 286 1810 mouseest@w lone is ava	Mouse EST Proje EST Project sity School of Parkway, Box 85 atson.wustl.edu ilable royalty-	MedicineP 01, St. Louis free through	LLNL ; contac		
	MGI:39 Seq pr	4246 imer: -28m1	(info@image.lln 3 rev2 ET from ence stop: 461.		rther informa	tion.	
source	e	1470 /organisi/strain= /db_xref /clone=" /clone_1 /sex="ma /tissue_ /dev_stai/lab_hos /note="V polylink was prim TGTTACCA 3']; dou (Pharmac and Eco provided	type="Thymus" ge="4 weeks" t="DH10B" ector: pT7T3D-P er; Site_1: Not ed with a Not I ATCTGAAGTGGGAGC ble-stranded cD ia), digested w RI sites of the by Dr. Bertran	ac (Pharmacia I; Site_2: E - oligo(dT) GGCCGCGTTTTTT NA was ligated ith Not I and modified pT7' d Jordan. Lib:	co RI; 1st st primer [5' TTTTTTTTTTT d to Eco RI a cloned into T3 vector. RN rary went thr	rand cDNA TTTTTT daptors the Not I A ough two	
BASE COUNT	1 2	Soares a	f normalization nd M.Fatima Bon		eciaciea by B	CIICO	

BASE COUNT 130 a 101 c 92 g 147 t

ORIGIN

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1 gtgatggtgc tatttattat tgcctttgag ccatctaccc tgattatcct atacacattt 61 cttgtgctgt cataaaaata agatacagta actgcatgct tgctggtggg tacccagttc 121 ttctttgtgt tcgggtcaat ctggaagaca tgagctcgag tgctgaagat aggttgctcc 181 cccattttgc ccaatgaaaa cttgctctga agtttctgct cttatgctac agaaataact 241 tccaaggaat tttctccgtc tgttattttg cagttgcttt tccacccccg agatcttttc 301 cggagcaact caacaaggca gacaattcag ttagttcttg tcggtaaaat gatttcactc 361 gcgctgacac caaaatgttc aaacatttgg aatttgaaaa ggttgcggg agaatatagc 421 acttaactat tccacaaaat taagcgtacg tgtagaaaaa acgcaaaagc
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